Amendments to the Specification:

Please replace the paragraph beginning at page 13, line 25, with the following:

--Amino acids that may be used to link the S3 multimer of interest to the remainder of the polypeptide include aspartic acid-proline, asparagine-glycine, methionine, cysteine, lysine-proline, arginine-proline, isoleucine-glutamic acid-glycine-arginine (SEQ ID NO:10), and the like. Cleavage may be effected by exposure to the appropriate chemical reagent or cleaving enzyme.--

Please replace the paragraph beginning at page 14, line 24, with the following:

--Examples of enzymatic agents include proteases, such as collagenase, which in some cases ~~recognizes~~ recognize the amino acid sequence NH_2--Pro--X--Gly--Pro--COOH, wherein X is an arbitrary amino acid residue, e. g. leucine; chymosin (rennin), which cleaves the Met-Phe bond; kallikrein B, which cleaves on the carboxyl side of Arg in X--Phe--Arg--Y; enterokinase, which recognizes the sequence X--$(Asp)_n$--Lys--Y, wherein n=2-4 (SEQ ID NO:11), and cleaves it on the carboxyl side of Lys; and thrombin which cleaves at specific arginyl bonds. Examples of chemical agents include cyanogen bromide (CNBr), which cleaves after Met; hydroxylamine, which cleaves the Asn-Z bond, wherein Z may be Gly, Leu or Ala; formic acid, which in high concentration (about 70%) specifically cleaves Asp-Pro. Thus, if the desired portion does not contain any methionine sequences, the cleavage site may be a methionine group which can be selectively cleaved by cyanogen bromide.--

Please cancel the present "SEQUENCE LISTING", pages 1/12-12/12, submitted for parent Application No. PCT/SG04/000194, and insert therefor the accompanying paper copy of the Substitute Sequence Listing, page numbers 1 to 12, at the end of the application.